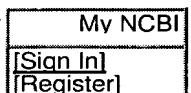


# **EXHIBIT**

**1**



PubMed Nucleotide Protein Genome Structure PMC Taxonomy OMIM Books

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Range: from  to  Features:  SNP graph  CDD  MGC  HPRD  STS  tRNA  Refre:

**1: 721P.** Reports Chain , H-Ras P2...[gi:494925]

BLink, Conserved Domains, Links

LOCUS 721P 166 aa linear PRI 07-OCT-1998  
DEFINITION H-Ras P21 Protein Mutant With Gln 61 Replaced By Leu (Q61L) Complex

With Guanosine-5'-[b,G-Imido] Triphosphate.

ACCESSION 721P

VERSION 721P GI:494925

DBSOURCE pdb: molecule 721P, chain 32, release Jun 6, 1991;

deposition: Jun 6, 1991;

class: Oncogene Protein;

source: Human (Homo Sapiens) Cellular Harvey-Ras Gene Truncated And Expressed In (Escherichia Coli);

Exp. method: X-Ray Diffraction.

KEYWORDS .

SOURCE Homo sapiens (human)

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (residues 1 to 166)

AUTHORS Pai,E.F., Krengel,U., Petsko,G.A., Goody,R.S., Kabsch,W. and Wittinghofer,A.

TITLE Refined crystal structure of the triphosphate conformation of H-ras p21 at 1.35 Å resolution: implications for the mechanism of GTP hydrolysis

JOURNAL EMBO J. 9 (8), 2351-2359 (1990)

PUBMED 2196171

REFERENCE 2 (residues 1 to 166)

AUTHORS Krengel,U., Schlichting,L., Scherer,A., Schumann,R., Frech,M., John,J., Kabsch,W., Pai,E.F. and Wittinghofer,A.

TITLE Three-dimensional structures of H-ras p21 mutants: molecular basis for their inability to function as signal switch molecules

JOURNAL Cell 62 (3), 539-548 (1990)

PUBMED 2199064

REFERENCE 3 (residues 1 to 166)

AUTHORS Krengel,U., Scherer,A., Kabsch,W., Wittinghofer,A. and Pai,E.F.

TITLE Direct Submission

JOURNAL Submitted (06-JUN-1991)

COMMENT Revision History:

JAN 31 94 Initial Entry.

FEATURES Location/Qualifiers

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Plaint according to the invention!

P. 52 / l. 24

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Display **GenBank** Show **5** Send toRange: from  to  Reverse complemented strand Features:  SNP graph  CDD  MGC  HPRD  STS  tRNA**1: AB006590.** Reports **Homo sapiens mRNA...[gi:2911151]**

Links

**LOCUS** AB006590 1740 bp mRNA linear PRI 05-FEB-1999  
**DEFINITION** Homo sapiens mRNA for estrogen receptor beta, complete cds.  
**ACCESSION** AB006590  
**VERSION** AB006590.1 GI:2911151  
**KEYWORDS** estrogen receptor beta.  
**SOURCE** Homo sapiens (human)  
**ORGANISM** Homo sapiens  
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;  
Hominidae; Homo.  
**REFERENCE** 1 (sites)  
**AUTHORS** Ogawa,S., Inoue,S., Watanabe,T., Hiroi,H., Orimo,A., Hosoi,T.,  
Ouchi,Y. and Muramatsu,M.  
**TITLE** The complete primary structure of human estrogen receptor beta (hER  
beta) and its heterodimerization with ER alpha in vivo and in vitro  
**JOURNAL** Biochem. Biophys. Res. Commun. 243 (1), 122-126 (1998)  
**PUBMED** 9473491  
**REFERENCE** 2 (bases 1 to 1740)  
**AUTHORS** Ogawa,S.  
**TITLE** Direct Submission  
**JOURNAL** Submitted (13-AUG-1997)-Sumito Ogawa, Saitama Medical School,  
Department of 2nd Biochemistry; 38 Morohongo, Moroyama, Iruma-gun,  
Saitama 350-0495, Japan (E-mail:sugawa@saitama-med.ac.jp,  
Tel:81-492-76-1490, Fax:81-492-94-9751)  
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## 1: 1204262A. Reports estrogen receptor...[gi:224957]

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LOCUS 1204262A 595 aa linear PRI 10-AUG-1994

DEFINITION estrogen receptor

ACCESSION 1204262A

VERSION 1204262A GI:224957

DBSOURCE prf: locus 1204262A;

state: fibrosarcoma;

taxonomy: Mammalia

KEYWORDS Estrogen Receptor; Human; cDNA Clone; Breast Cancer; Seq Determination; 6460bp; 595AAs; Expression in HeLa Cell; Binding of Estradiol; Seq Homol with erbA Protein; Hydropathy Plot.

SOURCE Homo sapiens (human)

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (residues 1 to 595)

AUTHORS Green,S., Walter,P., Kumar,V., Krust,A., Bornert,J.M., Argos,P. and Chambon,P.

TITLE Human oestrogen receptor cDNA: sequence, expression and homology to v-erb-A

JOURNAL Nature 320 (6058), 134-139 (1986)

PUBMED 3754034

COMMENT cDNA.

FEATURES Location/Qualifiers

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Range: from **begin** to **end** Reverse complemented strand Features:  SNP graph  CDD  MGC  HPRD  STS  tRNA

**1: X51416. Reports Human mRNA for st...[gi:36608]**

Links

**LOCUS** HSSTHOR 2402 bp mRNA linear PRI 18-APR-2005  
**DEFINITION** Human mRNA for steroid-hormone-receptor-hERR1.  
**ACCESSION** X51416 Y00290  
**VERSION** X51416.1 GI:36608  
**KEYWORDS** hormone receptor; receptor; steroid hormone receptor; transmembrane protein.  
**SOURCE** Homo sapiens (human)  
**ORGANISM** Homo sapiens  
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;  
Homidae; Homo.  
**REFERENCE** 1 (bases 1 to 2402)  
**AUTHORS** Giguere,V., Yang,N., Segui,P. and Evans,R.M.  
**TITLE** Identification of a new class of steroid hormone receptors  
**JOURNAL** Nature 331 (6151), 91-94 (1988)  
**PUBMED** 3267207  
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BLINK, Conserved Domains, Links

**LOCUS** 1JAI 166 aa linear PRI 15-DEC-1996  
**DEFINITION** H-Ras P21 Protein-Mutant G12p, Complexed With Guanosine-5'-[beta,Gamma-Methylene] Triphosphate And Manganese.  
**ACCESSION** 1JAI  
**VERSION** 1JAI GI:2392390  
**DBSOURCE** pdb: molecule 1JAI, chain 32, release Dec 15, 1996;  
deposition: Dec 15, 1996;  
class: Gtp-Binding;  
source: Mol\_id: 1; Organism\_scientific: Homo Sapiens;  
Organism\_common: Human; Gene: H-Ras-1; Expression\_system: Escherichia Coli; Expression\_system\_strain: Ck 600 K;  
Expression\_system\_vector: Ptac Ras;  
Exp. method: X-Ray Diffraction.  
**KEYWORDS**  
**SOURCE** Homo sapiens (human)  
**ORGANISM** Homo sapiens  
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;  
Hominidae; Homo.  
**REFERENCE**  
**AUTHORS** Seeburg,P.H., Colby,W.W., Capon,D.J., Goeddel,D.V. and Levinson,A.D.  
**TITLE** Biological properties of human c-Ha-ras1 genes mutated at codon 12  
**JOURNAL** Nature 312 (5989), 71-75 (1984) 6092966  
**PUBMED**  
**REFERENCE**  
**AUTHORS** Pai,E.F., Krengel,U., Petsko,G.A., Goody,R.S., Kabsch,W. and Wittinghofer,A.  
**TITLE** Refined crystal structure of the triphosphate conformation of H-ras p21 at 1.35 A resolution: implications for the mechanism of GTP hydrolysis  
**JOURNAL** EMBO J. 9 (8), 2351-2359 (1990) 2196171  
**PUBMED**  
**REFERENCE**  
**AUTHORS** Schweins,T., Scheffzek,K., Assheuer,R. and Wittinghofer,A.  
**TITLE** The role of the metal ion in the p21ras catalysed GTP-hydrolysis: Mn<sup>2+</sup> versus Mg<sup>2+</sup>  
**JOURNAL** J. Mol. Biol. 266 (4), 847-856 (1997) 9102473  
**PUBMED**  
**REFERENCE**  
**AUTHORS** Schweins,T., Scheffzek,K., Assheuer,R. and Wittinghofer,A.  
**TITLE** Direct Submission  
**JOURNAL** Submitted (15-DEC-1996)  
**COMMENT** Revision History:  
JUL 23 97 Initial Entry.  
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### **F 1: AAC13246. Reports ras p21 [Canis fa...[gi:3043763]**

LOCUS AAC13246 83 aa linear MAM 25-APR-2000  
 DEFINITION ras p21 [Canis familiaris].  
 ACCESSION AAC13246  
 VERSION AAC13246.1 GI:3043763  
 DBSOURCE locus CFU62092 accession U62092.3  
 KEYWORDS .  
 SOURCE Canis familiaris (dog)  
 ORGANISM Canis familiaris  
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 Mammalia; Eutheria; Laurasiatheria; Carnivora; Fissipedia; Canidae;  
 Canis.  
 REFERENCE 1 (residues 1 to 83)  
 AUTHORS Watzinger,F., Mayr,B., Haring,E. and Lion,T.  
 TITLE High sequence similarity within ras exons 1 and 2 in different  
 mammalian species and phylogenetic divergence of the ras gene  
 family  
 JOURNAL Mamm. Genome 9 (3), 214-219 (1998) 1  
 PUBMED 9501305  
 REFERENCE 2 (residues 1 to 83)  
 AUTHORS Watzinger,F.  
 TITLE Direct Submission  
 JOURNAL Submitted (25-JUN-1996) Children's Cancer Research Institute,  
 Kinderspitalgasse 6, Vienna A-1090, Austria  
 REFERENCE 3 (residues 1 to 83)  
 AUTHORS Watzinger,F.  
 TITLE Direct Submission  
 JOURNAL Submitted (23-NOV-1999) Children's Cancer Research Institute,  
 Kinderspitalgasse 6, Vienna A-1090, Austria  
 REMARK Sequence update by submitter  
 REFERENCE 4 (residues 1 to 83)  
 AUTHORS Watzinger,F.  
 TITLE Direct Submission  
 JOURNAL Submitted (25-APR-2000) Children's Cancer Research Institute,  
 Kinderspitalgasse 6, Vienna A-1090, Austria  
 REMARK Sequence update by submitter  
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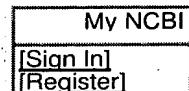
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### 1: AAC13249. Reports ras p21 [Felis ca...[gi:3043769]

[BLINK](#), [Conserved Domains](#), [Links](#)

LOCUS AAC13249 83 aa linear MAM 10-APR-1998  
 DEFINITION ras p21 [Felis catus].  
 ACCESSION AAC13249  
 VERSION AAC13249.1 GI:3043769  
 DBSOURCE locus FCU62088 accession U62088.1  
 KEYWORDS .  
 SOURCE Felis catus (cat)  
 ORGANISM Felis catus  
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 Mammalia; Eutheria; Laurasiatheria; Carnivora; Fissipedia; Felidae;  
 Felinae; Felis.  
 REFERENCE 1 (residues 1 to 83)  
 AUTHORS Watzinger,F., Mayr,B., Haring,E. and Lion,T.  
 TITLE High sequence similarity within ras exons 1 and 2 in different  
 mammalian species and phylogenetic divergence of the ras gene  
 family  
 JOURNAL Mamm. Genome 9 (3), 214-219 (1998)  
 PUBMED 9501305  
 REFERENCE 2 (residues 1 to 83)  
 AUTHORS Watzinger,F.  
 TITLE Direct Submission  
 JOURNAL Submitted (25-JUN-1996) Children's Cancer Research Institute,  
 Kinderspitalgasse 6, Vienna A-1090, Austria  
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## 1: 621P. Reports Chain , H-Ras P2...[gi:494922]

BLINK,  Conserved Domains,  Links

LOCUS 621P 166 aa linear PRI 07-OCT-1998  
DEFINITION H-Ras P21 Protein Mutant With Gln 61 Replaced By His (Q61h) Complex With Guanosine-5'-[b,G-Imido]Triphosphate.

ACCESSION 621P  
VERSION 621P GI:494922  
DBSOURCE pdb: molecule 621P, chain 32, release Jun 6, 1991;  
deposition: Jun 6, 1991;  
class: Oncogene Protein;  
source: Human (Homo Sapiens) Cellular Harvey-Ras Gene Truncated And Expressed In (Escherichia Coli);  
Exp. method: X-Ray Diffraction.

KEYWORDS  
SOURCE  
ORGANISM Homo sapiens (human)  
Homo sapiens  
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (residues 1 to 166)  
AUTHORS Pai,E.F., Krengel,U., Petsko,G.A., Goody,R.S., Kabsch,W. and Wittinghofer,A.  
TITLE Refined crystal structure of the triphosphate conformation of H-ras p21 at 1.35 Å resolution: implications for the mechanism of GTP hydrolysis

JOURNAL EMBO J. 9 (8), 2351-2359 (1990)  
PUBMED 2196171

REFERENCE 2 (residues 1 to 166)  
AUTHORS Krengel,U., Schlichting,L., Scherer,A., Schumann,R., Frech,M., John,J., Kabsch,W., Pai,E.F. and Wittinghofer,A.  
TITLE Three-dimensional structures of H-ras p21 mutants: molecular basis for their inability to function as signal switch molecules

JOURNAL Cell 62 (3), 539-548 (1990)  
PUBMED 2199064

REFERENCE 3 (residues 1 to 166)  
AUTHORS Krengel,U., Scherer,A., Kabsch,W., Wittinghofer,A. and Pai,E.F.

TITLE Direct Submission

JOURNAL Submitted (06-JUN-1991)

COMMENT Revision History:

JAN 31 94 Initial Entry.

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### 1: AAB02605. Reports c-Ha-ras1 p21 pro...[gi:190891]

LOCUS AAB02605 . . . . . 189 aa . . . . . linear PRI 13-FEB-2004

DEFINITION c-Ha-ras1 p21 protein [Homo sapiens]?

ACCESSION AAB02605 CAA23837

VERSION AAB02605.1 GI:190891

DBSOURCE locus HMRASH accession J00277.1

KEYWORDS

SOURCE Homo sapiens (human)

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 5 (residues 1 to 189)

AUTHORS Capon,D.J., Chen,E.Y., Levinson,A.D., Seeburg,P.H. and Goeddel,D.V.

TITLE Complete nucleotide sequences of the T24 human bladder carcinoma oncogene and its normal homologue

JOURNAL Nature 302 (5903), 33-37 (1983)

PUBMED 6298635

COMMENT On Feb 12, 2004 this sequence version replaced gi:35887.

The human genome contains a family of genes with homology to the Harvey murine sarcoma virus oncogene (c-Ha-ras). Two of these homologues are detectable by high stringency Southern hybridizations; the c-Ha-ras1 and c-Ha-ras2 genes, which occur on BamHI fragments of varying sizes from 3 to 9 kb because of allelic polymorphisms in the flanking regions.

Genomic mapping and nucleotide sequencing has shown that c-Ha-ras1 is the normal progenitor of the transforming gene found in several human tumor cell lines (T24 bladder carcinoma; EJ bladder carcinoma; Hs242 lung carcinoma; SK2 melanoma; and HS578T mammary carcinosarcoma). The only difference within the coding exons of the c-Ha-ras1 proto-oncogene and the oncogene of the T24 and EJ cell lines is a 'g' to 't' transversion within codon 12 that results in the substitution of valine for glycine at this position in the p21 protein encoded by c-Ha-ras1. The mutation responsible for transforming ability of the p21 protein in the SK2 and Hs242 cell lines is an 'a' to 't' transversion within codon 61 that results in the substitution of leucine for glutamine at this position. The mutation responsible for transforming ability of the p21 protein in the HS578T cell line is a 'g' to 'a' transition within codon 12 that results in the substitution of Aspartic acid for glycine at this position [9]. Because this 'g' to 'a' transition abolishes a MspI/HpaII site within the transformed allele, [9] was able to determine that both of the H-ras1 alleles found in normal cells from the same individual from which the HS578T cells were obtained, had the normal sequence at this position.

A region of repeated DNA consisting of the 28bp consensus sequence 'cactcccccttctccaggggacgcca' begins at position 4755. The repeat occurs 29 times in the plasmid used for this sequence but may occur more times in the native DNA. This region is known to be unessential for transforming activity [5].

[5] constructs a chimeric SV40 early promoter/human c-Ha-ras1 plasmid to demonstrate that upon transfection transforming activity is unaffected. [2], [5] and [7] discuss the extensive homology with the retroviral onc genes (v-has, v-bas, v-ha-ras). Complete source information:

Human genomic DNA [3], [1], [5]; human bladder carcinoma cell line T24 DNA [3], [2], [5], [7]; cDNA to mRNA, clones RS-3, RS-4 and RS-6 [4]; human bladder carcinoma cell line EJ DNA [1]; human lung carcinoma cell line Hs242 DNA [6]; human melanoma cell line SK2 DNA, clone lambda-SK2-T2 [8]; human mammary carcinosarcoma cell line SH578T DNA, clone lambda-HS578T [9], [10].

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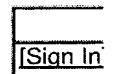
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Page 1 of 2 Next

Γ 1: 1WQ1G Reports

Chain G, Ras-Rasgap Complex

gi|3402130|pdb|1WQ1|G[3402130]

BLink, Conserved Domains, Links

Γ 2: 1WQ1R Reports

Chain R, Ras-Rasgap Complex

gi|3402129|pdb|1WQ1|R[3402129]

BLink, Conserved Domains, Links

Γ 3: 1A2B Reports

Chain , Human Rhoa Complexed With Gtp Analogue

gi|3318980|pdb|1A2B| [3318980]

BLink, Conserved Domains, Links

Γ 4: 2RGF Reports

Chain , Rbd Of Ral Guanosine-Nucleotide Exchange Factor (Protein),  
Nmr, 10 Structures

gi|2982004|pdb|2RGF| [2982004]

BLink, Conserved Domains, Links

Γ 5: 1JAI Reports

Chain , H-Ras P21 Protein Mutant G12p, Complexed With  
Guanosine-5'-[beta,Gamma-Methylene] Triphosphate And Manganese

gi|2392390|pdb|1JAI| [2392390]

BLink, Conserved Domains, Links

Γ 6: 1JAH Reports

Chain , H-Ras P21 Protein Mutant G12p, Complexed With  
Guanosine-5'-[beta,Gamma-Methylene] Triphosphate And Magnesium

gi|2392389|pdb|1JAH| [2392389]

BLink, Conserved Domains, Links

Γ 7: Reports

BAA20127

Rap 1B [Rattus norvegicus]

gi|2116982|dbj|BAA20127.1|[2116982]

BLink, Conserved Domains, Links

Γ 8: Reports

BAA20126

Rap 1A [Rattus norvegicus]

gi|2116980|dbj|BAA20126.1|[2116980]

BLink, Conserved Domains, Links

Γ 9: Reports

AAC52724

RalGDS-like factor

gi|1354501|gb|AAC52724.1|[1354501]

BLink, Conserved Domains, Links

Γ 10: 1AGP Reports

Chain , C-H-Ras P21 Protein Mutant With Gly 12 Replaced By Asp

BLink, Conserved Domains, Links

(G12d) Complexed With Guanosine-5'-[b,G-Imido] Triphosphate  
gi|515076|pdb|1AGP| [515076]

Γ 11: 1GNP Reports BLink, Conserved Domains, Links  
Chain , C-H-Ras P21 Protein Complexed With 3'-O-(N-Methyl-Anthraniloyl-2'-Deoxyguanosine-5'-[beta,Gamma-Imido]- Triphosphate (Residues 1 - 166)  
gi|1127267|pdb|1GNP| [1127267]

Γ 12: 1PLL Reports BLink, Conserved Domains, Links  
Chain , C-H-Ras P21 Protein Mutant With Gly 12 Replaced By Pro (G12p) Complexed With Guanosine-Diphosphate  
gi|576244|pdb|1PLL| [576244]

Γ 13: 1PLK Reports BLink, Conserved Domains, Links  
Chain , C-H-Ras P21 Protein Mutant With Gly 12 Replaced By Pro (G12p) Complexed With Guanosine-Triphosphate  
gi|576243|pdb|1PLK| [576243]

Γ 14: 1PLJ Reports BLink, Conserved Domains, Links  
Chain , C-H-Ras P21 Protein Mutant With Gly 12 Replaced By Pro (G12p) Complexed With P3-1-(2-Nitrophenyl)ethyl-Guanosine-5'-(B,G-Imido)-Triphosphate  
gi|576242|pdb|1PLJ| [576242]

Γ 15: 821P Reports BLink, Conserved Domains, Links  
Chain , C-H-Ras P21 Protein (Residues 1 - 166) Mutant With Gly 12 Replaced By Pro (G12p) Complex With Guanosine-5'-[b,G-Imido] Triphosphate  
gi|494936|pdb|821P| [494936]

Γ 16: 721P Reports BLink, Conserved Domains, Links  
Chain , H-Ras P21 Protein Mutant With Gln 61 Replaced By Leu (Q61l) Complex With Guanosine-5'-[b,G-Imido] Triphosphate  
gi|494925|pdb|721P| [494925]

Γ 17: 621P Reports BLink, Conserved Domains, Links  
Chain , H-Ras P21 Protein Mutant With Gln 61 Replaced By His (Q61h) Complex With Guanosine-5'-[b,G-Imido] Triphosphate  
gi|494922|pdb|621P| [494922]

Γ 18: 521P Reports BLink, Conserved Domains, Links  
Chain , H-Ras P21 Protein Mutant With Gly 12 Replaced By Val (G12v) Complex With Guanosine Triphosphate  
gi|494910|pdb|521P| [494910]

Γ 19: 421P Reports BLink, Conserved Domains, Links  
Chain , H-Ras P21 Protein Mutant With Gly 12 Replaced By Arg (G12r) Complex With Guanosine-5'-[b,G-Imido] Triphosphate  
gi|494886|pdb|421P| [494886]

Γ 20: 221P Reports BLink, Conserved Domains, Links  
Chain , H-Ras P21 Protein Mutant With Asp 38 Replaced By Glu (D38e) Complex With Guanosine-5'-[b,G-Imido] Triphosphate  
gi|494721|pdb|221P| [494721]